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(FILE 'HOME' ENTERED AT 15:50:06 ON 16 JAN 2009)

FILE 'MEDLINE, EMBASE, BIOSIS, SCISEARCH, DISSABS, TOXCENTER' ENTERED AT
15:51:09 ON 16 JAN 2009

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L1      2729 S (HEINRICH M?/AU) OR (CORLESS C?/AU)
L2     10611 S (FLETCHER J?/AU) OR (DEMETRI G?/AU)
L3     13149 S L1 OR L2
L4      218 S L3 AND (PDGF?)
L5       0 S L3 AND CD140?
L6     180 S L3 AND (PLATELET(1A)DERIVED)
L7     264 S L4 OR L6
L8     207 S L7 AND MUTAT?
L9     178 S L8 AND (TUMOR? OR CANCER? OR CARCIN?)
L10    109 S L8 AND NEOPLAS?
L11    186 S L9 OR L10
L12    456 S L3 AND GIST
L13    128 S L4 AND GIST
L14     78 DUP REM L11 (108 DUPLICATES REMOVED)
L15     7 S L14 AND PY<=2002
L16    199 DUP REM L12 (257 DUPLICATES REMOVED)
L17     26 S L16 AND PY<=2002
L18     31 S L15 OR L17
L19     31 DUP REM L18 (0 DUPLICATES REMOVED)
L20    2371 S (PDGFRA)
L21     99 S (PDGFR-A)
L22    2344 S (PDGFR(2A)ALPHA)
L23     327 S (PDGF-R(2A)ALPHA)
L24    2748 S (PDGF(2A)RECEPTOR(2A)ALPHA)
L25    4759 S (PLATELET(2A)DERIVED(5A)RECEPTOR(10A)ALPHA)
L26     23 S CD140A
L27    8628 S L20 OR L21 OR L22 OR L23 OR L24 OR L25 OR L26
L28    2386 S L27 AND MUTAT?
L29     0 S L27 AND (2072 OR 2073 OR 2074 OR 2075)
L30     1 S L27 AND (2076 OR 2077 OR 2078 OR 2079 OR 2080)
L31     0 S L27 AND (2081 OR 2082 OR 2083 OR 2084 OR 2085)
L32     0 S L27 AND (2086 OR 2087 OR 2088 OR 2089 OR 2090)
L33     0 S L27 AND (2091 OR 2092 OR 2093 OR 2094 OR 2095)
L34     0 S L27 AND (2096 OR 2097 OR 2098 OR 2099 OR 2100)
L35     0 S L27 AND (2101 OR 2102 OR 2103 OR 2104 OR 2105 OR 2106 OR 2107)
L36     0 S L27 AND (2916 OR 2917 OR 2918 OR 2919 OR 2920 OR 2921 OR 2922)
L37     0 S L27 AND (2923 OR 2924 OR 2925 OR 2926 OR 2927 OR 2928 OR 2929)
L38     0 S L27 AND (2930 OR 2931 OR 2932 OR 2933 OR 2934 OR 2935 OR 2936)
L39     39 S L27 AND (560 OR 561 OR 562 OR 563 OR 564 OR 565)
L40    156 S L27 AND (566 OR 567 OR 568 OR 569 OR 570 OR 571)
L41    186 S L39 OR L40
L42    117 DUP REM L41 (69 DUPLICATES REMOVED)
L43     48 S L27 AND (841 OR 842 OR 843 OR 844 OR 845 OR 846 OR 847 OR 848)
L44    161 S L42 OR L43
L45    127 DUP REM L44 (34 DUPLICATES REMOVED)
L46     70 S L45 AND MUTAT?
L47     27 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(560 OR 561 OR 562
L48     11 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(566 OR 567 OR 568
L49     37 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(841 OR 842 OR 843
L50     16 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(845 OR 846 OR 847
L51     60 S L47 OR L48 OR L49 OR L50
L52    20 DUP REM L51 (40 DUPLICATES REMOVED)
L53     0 S L52 AND PY<=2002
L54     2 S L52 AND PY<=2003
L55    349 S L27 AND ((EXON(1A)10)OR(EXON(1A)12))
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L56      111 DUP REM L55 (238 DUPLICATES REMOVED)
L57      1 S L56 AND PY<=2002
L58     186 S L27 AND ((EXON(1A)18))
L59      67 DUP REM L58 (119 DUPLICATES REMOVED)
L60      0 S L59 AND PY<=2002
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=> log y